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Display Show Hide: ☐ sequence ☐ all but gene, CDS and mRNA

Range: from to ☐ Reverse complemented strand Features:

☐ 1: AY039589. Reports Arabidopsis thaliana...[gi:14517507]

Links

Comment Features Sequence

LOCUS AY039589 685 bp mRNA linear PLN 20-JUN-2001
DEFINITION Arabidopsis thaliana AT4g02380/T14P8_2 mRNA, complete cds.
ACCESSION AY039589
VERSION AY039589.1 GI:14517507
KEYWORDS FLI CDNA.
SOURCE Arabidopsis thaliana (thale cress)
ORGANISM Arabidopsis thaliana
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicotyledons;
rosids; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
REFERENCE 1 (bases 1 to 685)
AUTHORS Kim,C.J., Chen,H., Cheuk,R., Koesema,E., Meyers,M.C., Banh,J.,
Bowser,L., Carninci,P., Dale,J.M., Gibson,H.A., Goldsmith,A.D.,
Hayashizaki,Y., Ishida,J., Jiang,P.X., Jones,T., Kamiya,A.,
Karlin-Neumann,G., Kawai,J., Lam,B., Lee,J.M., Lin,J., Liu,S.X.,
Miranda,M., Narusaka,M., Nguyen,M., Onodera,C.S., Palm,C.J.,
Pham,P.K., Quach,H.L., Sakurai,T., Satou,M., Seki,M., Southwick,A.,
Tang,C.C., Toriumi,M., Yamada,K., Yu,G., Yu,S., Shinozaki,K.,
Davis,R.W., Theologis,A. and Ecker,J.R.
TITLE Arabidopsis cDNA clones
JOURNAL Unpublished
REFERENCE 2 (bases 1 to 685)
AUTHORS Kim,C.J., Chen,H., Cheuk,R., Koesema,E., Meyers,M.C., Banh,J.,
Bowser,L., Carninci,P., Dale,J.M., Gibson,H.A., Goldsmith,A.D.,
Hayashizaki,Y., Ishida,J., Jiang,P.X., Jones,T., Kamiya,A.,
Karlin-Neumann,G., Kawai,J., Lam,B., Lee,J.M., Lin,J., Liu,S.X.,
Miranda,M., Narusaka,M., Nguyen,M., Onodera,C.S., Palm,C.J.,
Pham,P.K., Quach,H.L., Sakurai,T., Satou,M., Seki,M., Southwick,A.,
Tang,C.C., Toriumi,M., Yamada,K., Yu,G., Yu,S., Shinozaki,K.,
Davis,R.W., Theologis,A. and Ecker,J.R.
TITLE Direct Submission
JOURNAL Submitted (04-JUN-2001) Salk Institute Genomic Analysis Laboratory
(SIGnAL), Plant Biology Laboratory, The Salk Institute for
Biological Studies, 10010 N. Torrey Pines Road, La Jolla, CA 92037,
USA
COMMENT RIKEN Genomic Sciences Center (GSC) members carried out the
collection and clustering of RAFL cDNAs (RAFL cDNA : 'RIKEN
Arabidopsis Full-Length cDNA') : Seki,M., Narusaka,M., Ishida,J.,
Satou,M., Kamiya,A., Sakurai,T., Carninci,P., Kawai,J.,
Hayashizaki,Y. and Shinozaki,K.

The Salk, Stanford, PGEC (SSP) Consortium members carried out the
sequencing and annotation of the RAFL cDNAs: Shinn,P., Chen,H.,

Cheuk,R., Kim,C.J., Koesema,E., Meyers,M.C., Banh,J. Bowser,L., Dale,J.M., Gibson,H.A., Goldsmith,A.D., Jiang,P.X., Jones,T., Karlin-Neumann,G., Lam,B., Lee,J.M., Lin,J., Liu,S.X., Miranda,M., Nguyen,M., Onodera,C.S., Palm,C.J., Pham,P.K., Quach,H.L., Southwick,A., Tang,C.C., Toriumi,M., Yamada,K., Yu,G., Yu,S., Davis,R.W., Theologis,A., and Ecker,J.R.

Shinn,P. (SSP/Salk) and Seki,M. (RIKEN GSC) contributed equally to this work. Shinozaki,K. (RIKEN GSC) and Ecker,J.R. (SSP/Salk) contributed equally to this work as PIs.

FEATURES
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3'UTR 484..685

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661 ttcttataaa accatttcag aactt
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Aug 28 2007 16:53:42